

RAW SEQUENCE LISTING ERROR REPORT

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Application Serial Number: 10/009, 980 A
Source: PC1/10
Date Processed by STIC: 11/18/2002

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FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216. PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax) PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

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- 1. EFS-Bio (http://www.uspto.gov/ebc/efs/downloads/documents.htm, EFS Submission User Manual ePAVE)
- 2. U.S. Postal Service: U.S. Patent and Trademark Office, Box Sequence, P.O. Box 2327, Arlington, VA 22202
- 3. Hand Carry directly to:

U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name, Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202

U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

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Revised 01/29/2002



PCT10

RAW SEQUENCE LISTING

DATE: 11/22/2002

PATENT APPLICATION: US/10/009,980A

TIME: 09:23:47

Input Set : N:\Crf4\11182002\J009980.raw Output Set: N:\CRF4\11222002\J009980A.raw

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2 <120> TITLE OF INVENTION: MOLECUTAR METHODS FOR DETECTING GUAR GUM ADDITIONS
3 TO LOCUST BEAN GUM LE Misspelled - Please Correct 4 <130> FILE REFERENCE: PATENT APPLICATION PCT/ES01/00079 Ly MOC
                                                              L) MOLECULAR.
5 <140> CURRENT APPLICATION NUMBER: US/10/009,980A
6 <141> CURRENT FILING DATE: 2002-07-29
                                                                         pp. 1-3.
7 <150> PRIOR APPLICATION NUMBER: ES200000560
8 <151> PRIOR FILING DATE: 2000-03-08
9 <160> NUMBER OF SEO ID NOS: 10
10 <170> SOFTWARE: PatentIn Ver. 2.1
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ERRORED SEQUENCES

Does Not Comply Corrected Diskette Needed Only Errored page

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E--> 140 delete

See p-2. for More Errors.

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/009,980A

DATE: 11/22/2002 TIME: 09:23:48

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L:80 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:180 L:81 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:240 L:139 M:254 E: No. of Bases conflict, this line has no nucleotides.

M:254 Repeated in SeqNo=10

L:140 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2 L:140 M:252 E: No. of Seq. differs, <211> LENGTH:Input:410 Found:411 SEQ:10